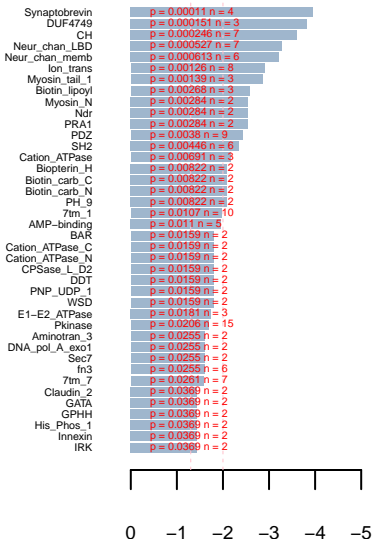


FM_Oe domain



log(p)

n = 578 genes | p adjust BH FALSE